JICSCI803 Algorithms and Data Structures March to June 2019

Highlights of Lecture 12

Parallel Algorithm
More Graph Algorithm

Parallel Algorithms

- Everything we have looked at so far has been based on one key assumption:
- Each instruction in an algorithm is performed one-at-a-time.
- In other words, all of the algorithms we have looked at are sequential.

Types of Parallelism

- We can classify parallel architectures by looking at the number of different instruction and data streams available in the architecture.
 - SIMD (Single instruction, Multiple data) -Array/Vector processors.
 - Each processor performs the same instruction on a different data element.
 - •MISD (Multiple instruction, Single data) –Pipeline processors.
 - Each processor performs a different instruction on a single data stream.
 - •MIMD (Multiple instruction, Multiple data) –high-level parallel processors.
 - Each processor operates fully independently of each of the others.

Types of Parallelism

– Consider the following code:

```
for i = 1, to 10 do
   if i is odd then
    x[i] = i
   else
   x[i] = 2 * I
```

-How would we code this for parallel execution?

MIMD

```
for i = 1, to 10 do in parallel
   if i is odd then
      x[i] = i
   else
      x[i] = 2 * I
```

- In this case each processing element performs one of the two possible calculations x[i] = i or x[i] = 2 * i.
- This is possible because the MIMD architecture
 allows processors to execute different instructions at the same time.

SIMD

```
for i = 1, to 10 step 2 do in parallel
    x[i] = i
    for i = 2, to 10 step 2 do in parallel
    x[i] = 2 * i
```

- In this case each processing element performs the same calculation at each step first x[i] = i and then x[i] = 2 * i.
- This is necessary because the SIMD architecture requires processors to execute the same instructions at the same time.

```
– Consider the following procedure:
  program MISDsort
    get max
    repeat
      get n
       if n < 0 then
          put max
      elseif n > max then
          put max
          max = n
      else
          put n
   until n < 0
   put -1
```

- Let us feed it the sequence 5 7 9 2 3 4 1 6 –1
- What is the output?

input	max	output
5	5	

input	max	output
5	5	
7	7	5

input	max	output
5	5	
7	7	5
9	9	7

input	max	output
5	5	
7	7	5
9	9	7
2	9	2

input	max	output
5	5	
7	7	5
9	9	7
2	9	2
3	9	3

input	max	output
5	5	
7	7	5
9	9	7
2	9	2
3	9	3
4	9	4

input	max	output
5	5	
7	7	5
9	9	7
2	9	2
3	9	3
4	9	4
1	9	1

input	max	output
5	5	
7	7	5
9	9	7
2	9	2
3	9	3
4	9	4
1	9	1
6	9	6

input	max	output
5	5	
7	7	5
9	9	7
2	9	2
3	9	3
4	9	4
1	9	1
6	9	6
– 1	9	9

input	max	output
5	5	
7	7	5
9	9	7
2	9	2
3	9	3
4	9	4
1	9	1
6	9	6
– 1	9	9
		<u> </u>

Input 5 7 **9** 2 3 4 1 6 -1
Output 5 7 2 3 4 1 6 9-1

- What has this process done?
- It has delayed the output of the largest element until last.
- Now consider what happens if we pipe the output of such a program into another copy of the same program.

in	max	out	in	max	out
5	5				

in	max	out	in	max	out
5	5				
7	7	5	5	5	

in	max	out	in	max	out
5	5				
7	7	5	5	5	
9	9	7	7	7	5

in	max	out	in	max	out
5	5				
7	7	5	5	5	
9	9	7	7	7	5
2	9	2	2	7	2

in	max	out	in	max	out
5	5				
7	7	5	5	5	
9	9	7	7	7	5
2	9	2	2	7	2
3	9	3	3	7	3
					_

in	max	out	in	max	out
5	5				
7	7	5	5	5	
9	9	7	7	7	5
2	9	2	2	7	2
3	9	3	3	7	3
4	9	4	4	7	4

in	max	out	in	max	out
5	5				
7	7	5	5	5	
9	9	7	7	7	5
2	9	2	2	7	2
3	9	3	3	7	3
4	9	4	4	7	4
1	9	1	1	7	1

in	max	out	in	max	out
5	5				
7	7	5	5	5	
9	9	7	7	7	5
2	9	2	2	7	2
3	9	3	3	7	3
4	9	4	4	7	4
1	9	1	1	7	1
6	9	6	6	7	6

in	max	out	in	max	out
5	5				
7	7	5	5	5	
9	9	7	7	7	5
2	9	2	2	7	2
3	9	3	3	7	3
4	9	4	4	7	4
1	9	1	1	7	1
6	9	6	6	7	6
– 1	9	9	9	9	7

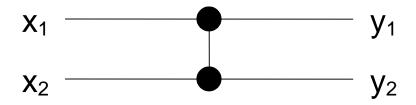
in	max	out	in	max	out
5	5				
7	7	5	5	5	
9	9	7	7	7	5
2	9	2	2	7	2
3	9	3	3	7	3
4	9	4	4	7	4
1	9	1	1	7	1
6	9	6	6	7	6
– 1	9	9	9	9	7
		– 1	– 1		9

in	max	out	in	max	out
5	5				
7	7	5	5	5	
9	9	7	7	7	5
2	9	2	2	7	2
3	9	3	3	7	3
4	9	4	4	7	4
1	9	1	1	7	1
6	9	6	6	7	6
– 1	9	9	9	9	7
		– 1	– 1		9
					– 1

Input 5 7 9 2 3 4 1 6 -1
Output 5 2 3 4 1 6 7 9 -1

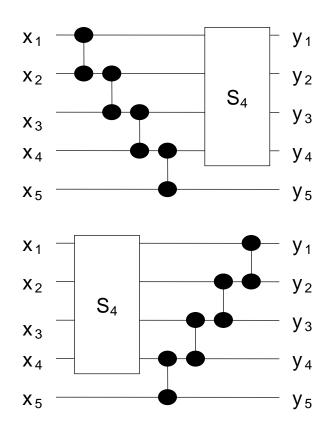
- Now the last two elements are in order.
- If we combine n-1 copies of the program in sequence we have a system to sort an n-element array.
- If we run the copies in parallel we have a sort algorithm which O(n).
- This mechanism of piping data through a series of parallel processes in sequence is the basis of MISD parallelism.

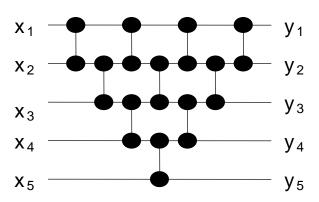
– Consider the following processing element:



- $-y_1 = \min(x_1, x_2)$
- $-y_2 = \max(x_1, x_2)$
- This is a comparator.
- Can we combine comparators in parallel to sort a set of numbers?

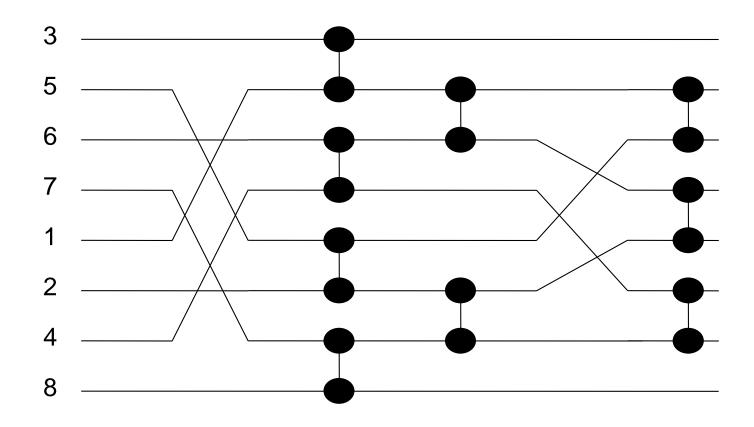
– If we have a circuit S_n which sorts an n-element set of numbers can we combine it with some additional comparators to produce S_{n+1} ?

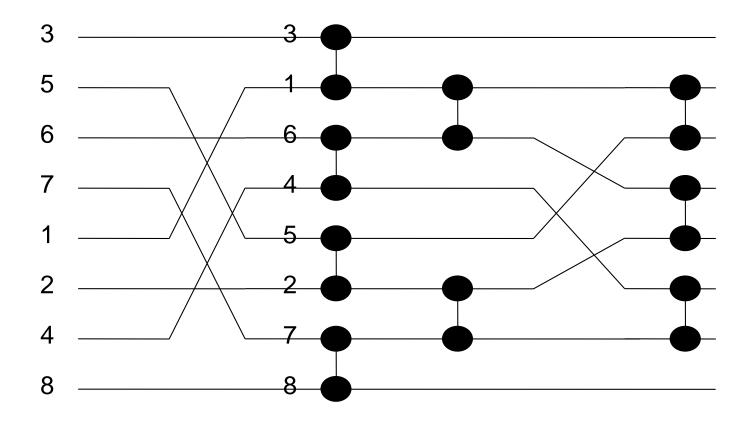


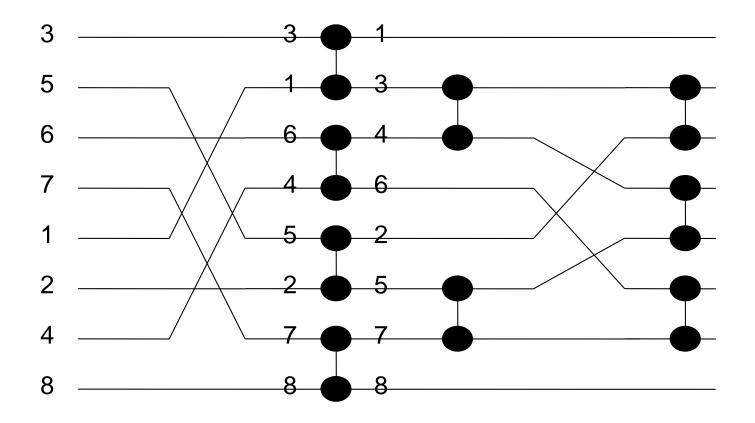


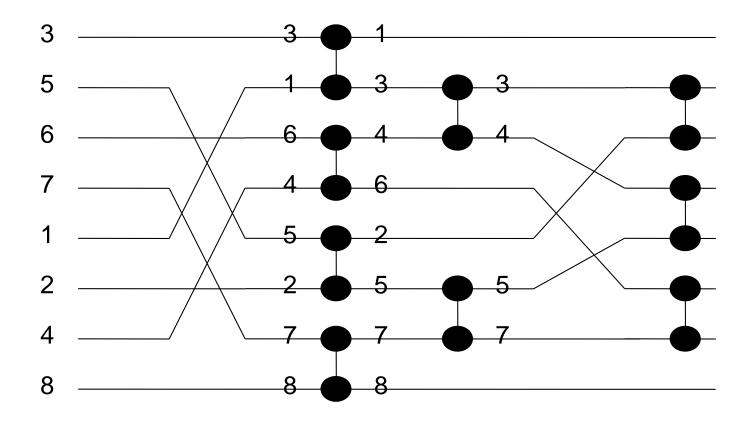
Parallel Sorting by Merging

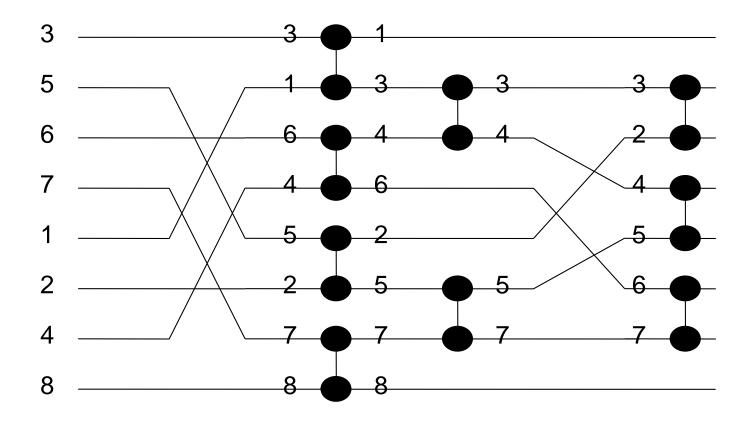
- Suppose we have two sorted streams, each of n elements.
- Can we use a method analogous to mergesort in parallel?

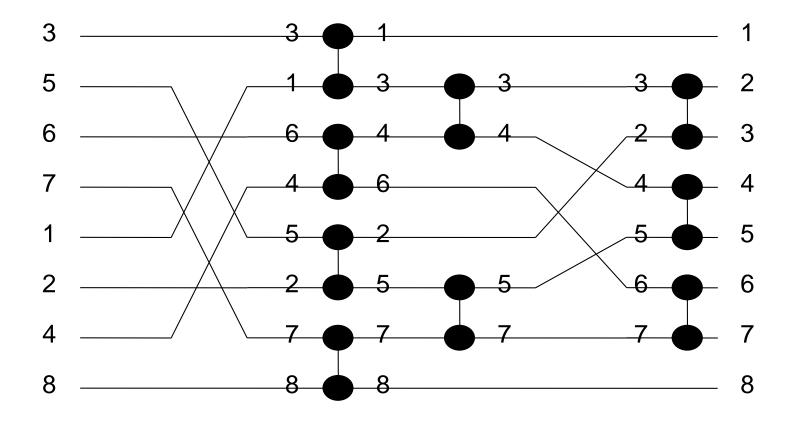






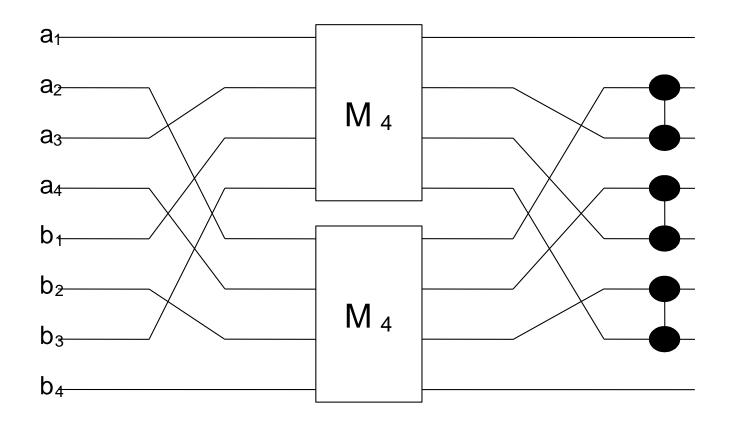




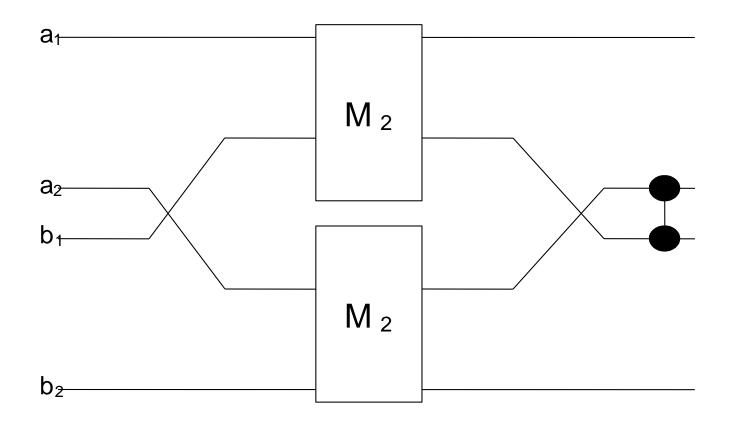


- Suppose we have two sorted streams, each of n elements.
- Can we use a method analogous to mergesort in parallel?
- Let us assume that we have a circuit that merges two n/2 element streams.

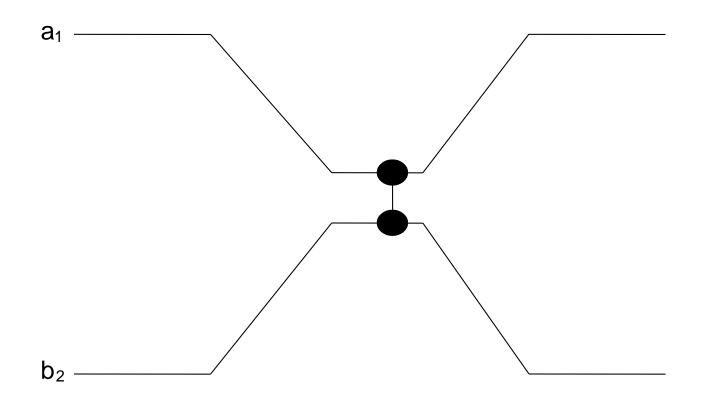
Parallel Sorting by Merging -M₈



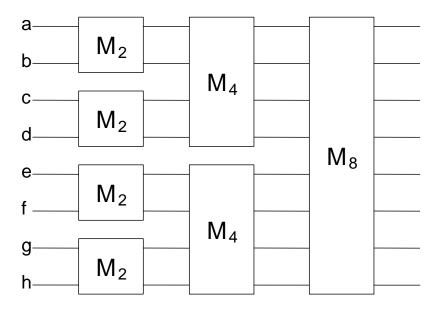
Parallel Sorting by Merging -M₄



Parallel Sorting by Merging –M₂



 We can combine these circuits to sort a parallel stream of numbers



More About Graphs

More About Graphs

- A great range of problems can be expressed in terms of graphs.
- We have already seen some of these:
 - Minimal spanning tree
 - Shortest path
- So far, all the algorithms we have examined have imposed an order on the nodes or the edges.
- This is not always necessary.

Graphs and Games

- Consider the following game:
 - Initially there is a heap of n matches between two players.
 - The first player may remove as many matches as she likes between 1 and n-1.
 - Thereafter each player can remove between 1 and twice the number their opponent just took.
 - The winner is the person who removes the last match.

The Match Game

- Consider the following situation:
- There is a pile of five matches in front of you and your opponent has just taken two matches. (the original pile has seven matches)
 - -Therefore, you can take 1, 2, 3 or 4 matches but not 5.
 - What should your next move be?
 - -If you take 2, 3, or 4 matches, your opponent can win next turn.
 - –So you should take 1 match.
 - But what about more complex positions?

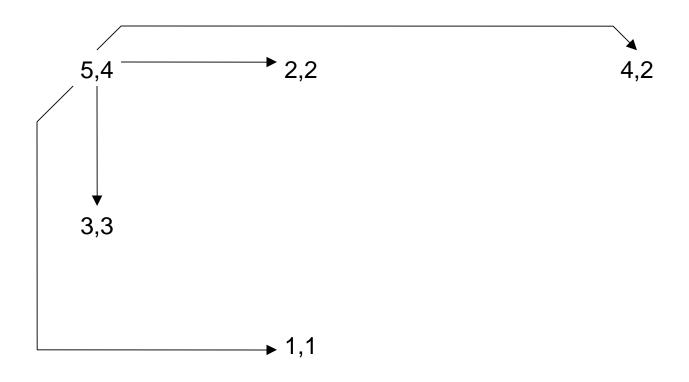
The Match Game as a graph

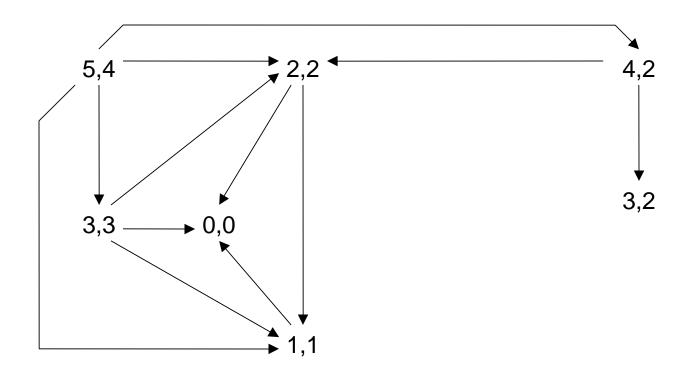
- We can represent positions in the match game as nodes on a graph.
- We can connect nodes to show sequences of possible moves.
- We label each node with two numbers:
 - The number of matches left at this stage of the game,
 - The number of matches which can be removed.
- Clearly node 0, 0 represents the end of the game.

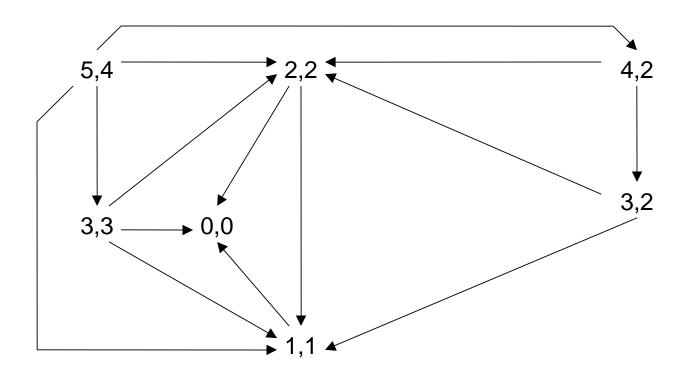
The Match Game as a graph

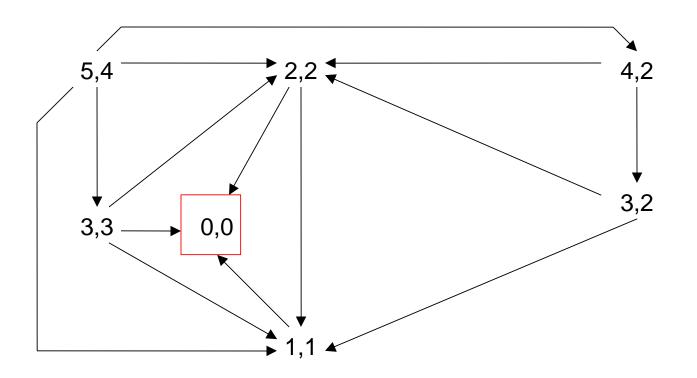
- Let losing nodes be squares and winning nodes be circles.
- Clearly 0, 0 is a losing node.
- 0, 0 can be reached from any node n, n.
- Any path leading to a winning node is a bad move.
- Any path leading to a losing node is a good move.
- Let us colour good moves in green.
- Let us colour bad moves in red.

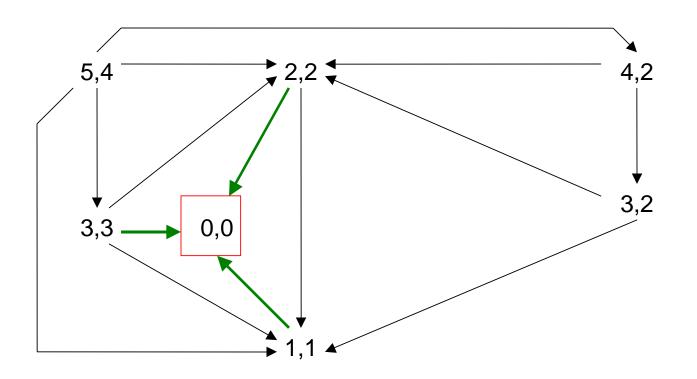
5,4

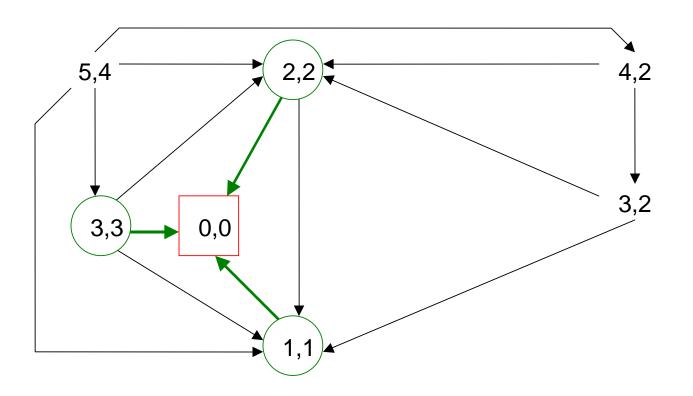


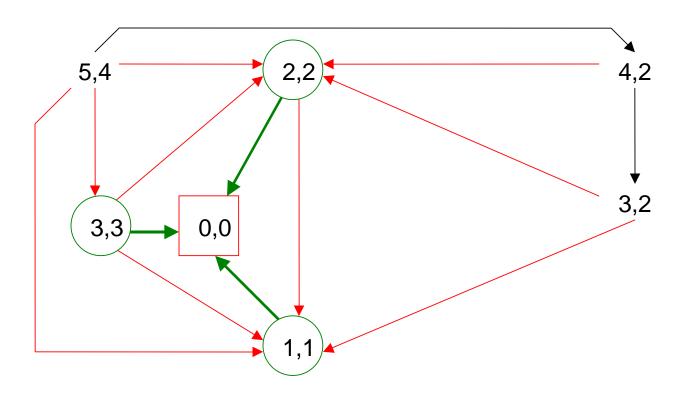


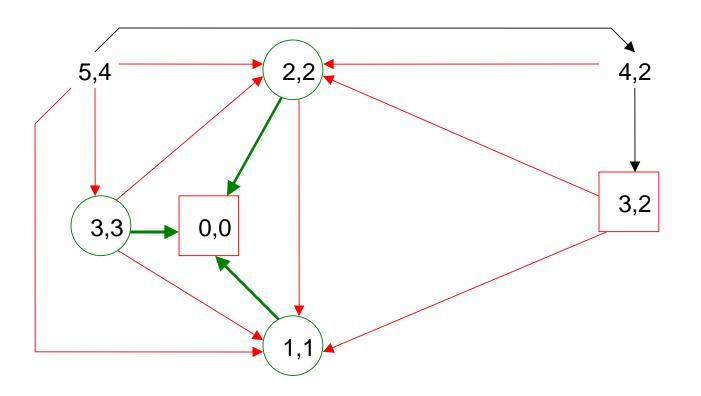


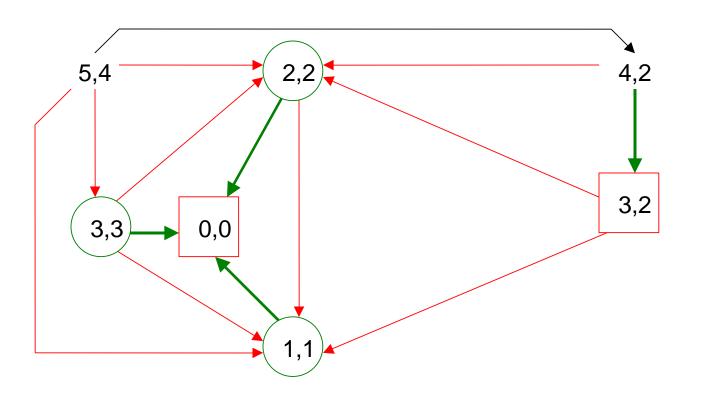


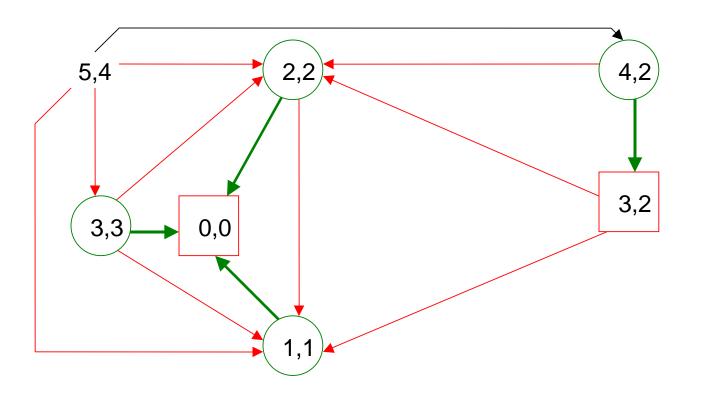


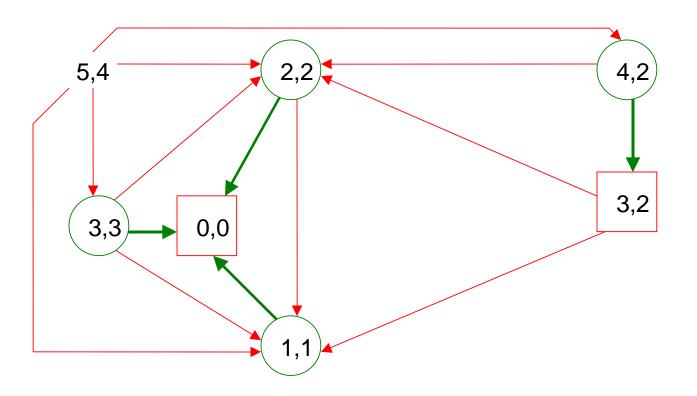


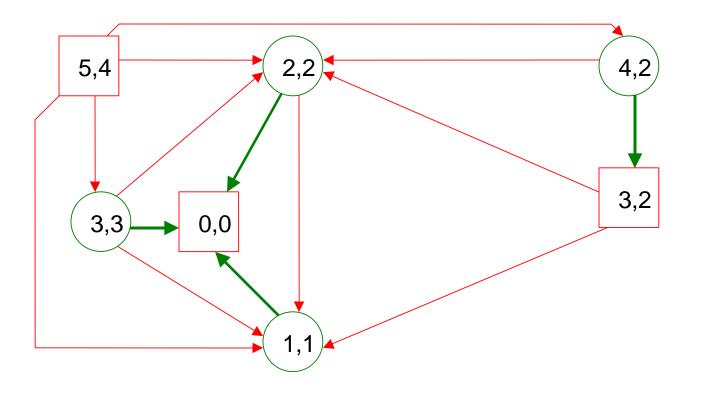


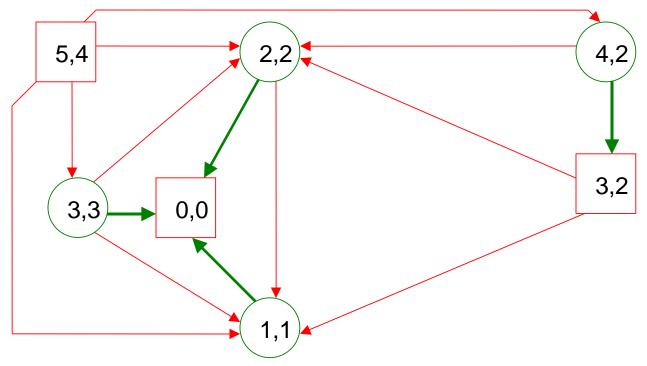












• So 5, 4 is a losing position.

Analysing the Match Game graph

- On a larger graph this process can be continued
- The rules can be summed up as follows:
 - A position is a winning position if at least one of its successors is a losing position.
 - A position is a losing position if all of its successors are winning positions.
- We can construct an algorithm that determines if we are at a winning position

```
function recwin(i, j)

// returns true if node i,j is winning

for k = 1 to j do

   if not recwin(i-k, min(2k, i-k)) then
    return true
```

return false

- This algorithm suffers the same problem as fibrec:
 - •Too may recursive calls
- -Can we fix this in some way?

- Can we fix this in some way?
- What if we remember the nodes we have already evaluated?
- Let us construct two arrays:
 - G[0..n, 0..n] where entry G[i, j] contains the value of the position i, j.
 - known[0..n, 0..n] where entry known[i, j] is true if G[i, j] has been evaluated.

```
G[0, 0] = false
 known[0,0] = true
  for i = 1 to n do
     for j = 1 to i do
        known[i, j] = false
function win(i, i)
     if known[i, j] then
         return G[i, j]
     known[i, j] = true
     for k = 1 to j do
         if not win(i-k, min(2k, i-k)) then
             G(i, j) = true
             return true
     G[i, j] = false
     return false
```

- This approach involves the initialisation of the array known[0..n,0..n]
- We can use virtual initialisation, described earlier, to eliminate this cost.
- This approach can be used to analyse positions in a number of games
- The basic process is the same

Analysing a general game

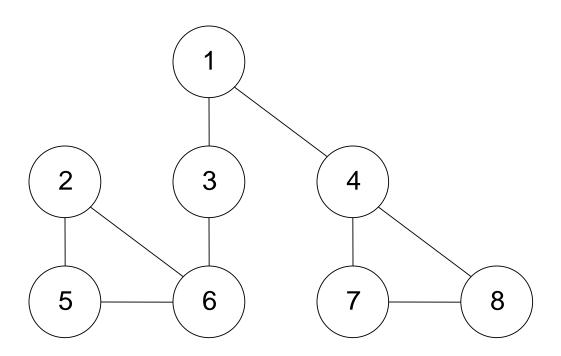
- Label the terminal position(s) win, lose, draw.
- A nonterminal position is a win if at least one of its successors is a losing position.
- A nonterminal position is a loss if all of its successors are winning positions.
- Any other nonterminal position is a draw.
- Once the graph is labelled, any position can be evaluated.

- Let $G = \langle N, A \rangle$ be an undirected graph.
- We wish to visit all the nodes of G.
- We wish to do this efficiently.
- Suppose we can mark a node to show it has been visited.
- Then we can construct a recursive algorithm to visit all of the nodes as follows.
- We will use a depth first algorithm to do this.

```
function dfvisit(G)
   for each v∈N do
      visited[v] = false
   for each v∈N do
      if not visited[v] then
            dfv[v]

function dfv[v]
   visited[v] = true
   for each node w adjacent to v do
      if not visited[w] then
            dfv[w]
```

– Consider the graph:



Traversing starting at node 1

dfv(1) initial call

dvf(3) recursive call

dfv(6) recursive call

dfv(2) recursive call

dfv(5) recursive call; progress blocked

dfv(4) an unvisited neighbour of node 1

dfv(7) recursive call

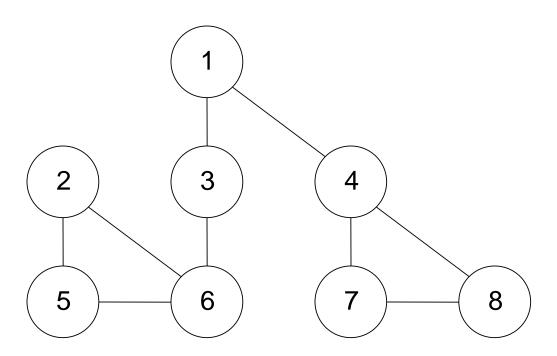
dfv(8) recursive call; progress blocked

• There are no more nodes to visit

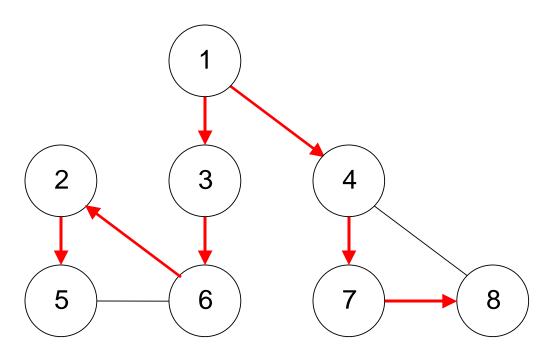
- Let G =⟨N, A⟩ have nnodes and aarcs (edges)
- How efficient is this algorithm?
 - Since each node is visited exactly once there are nealls to dfv.
 - At each node we examine connected nodes
 - This examination involves every edge being considered.
- The algorithm takes time in Θ (n) for procedure calls and a time in Θ (a) to inspect connected nodes.
- Overall the algorithm is in Θ (max(n, a))

- Depth first traversal associates a spanning tree T with the graph.
- The edges of T correspond to the edges used to traverse the graph.
- They are directed from the first node visited to the second.
- The starting point of the traversal becomes the root of the tree.
- If the graph is not connected, we produce a forest of scanning trees.

– Using our example graph:

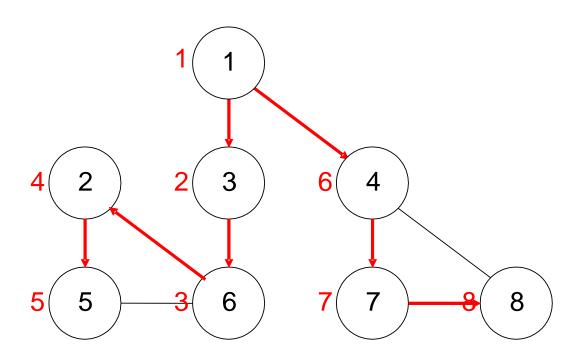


– Using our example graph, the spanning tree is:



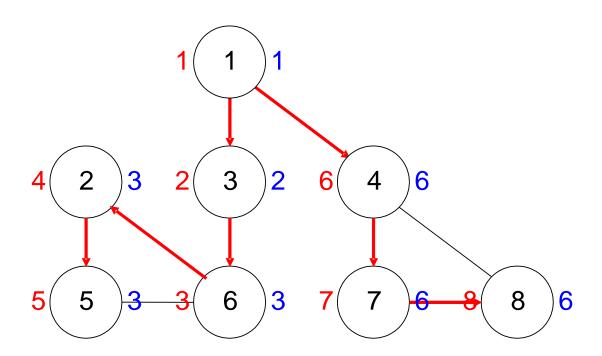
- A node v of a connected graph is an articulation point if the sub graph obtained by deleting node v and all its connected edges is no longer connected.
- Node 1 and node 3 are both articulation points in the example we have been using.
- Let us label the nodes of this graph to show the preorder traversal order for T.

– The preorder numbering is:



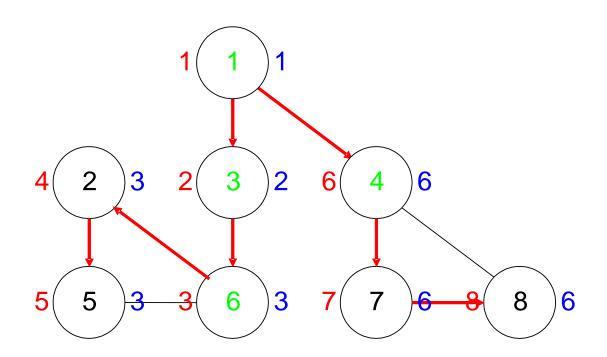
- Let us now number the graph again.
- In this instance the number has the following property:
 - Let w be the highest node that can be reached by following down zero or more arrows and then going along at most one non-arrow edge.
- Define prenum[w] is the serial number of w in the traversal process
- Define highest[v] = prenum[w]
- E.g. from node 7 we can reach node 4
 - Highest[7] = prenum[4] = 6

– The highest numbering is:



- Node w must be an ancestor of v
- Consider a node v which is not the root of T
- If v has no children it cannot be an articulation point
- Otherwise let x be a child of v
 - If highest[x] < prenum[v] then we can reach a node higher up the tree than node v from node x which means that node v need not be used to reach node x.
 - If highest[x]≥prenum[v] then node x can only be reached via node
- Node v is an articulation point if at least one of its children x satisfies highest[x] ≥ prenum[v]
- The root is an articulation point if it has more than one child.

Nodes 1, 3, 4 and 6 are articulation points.



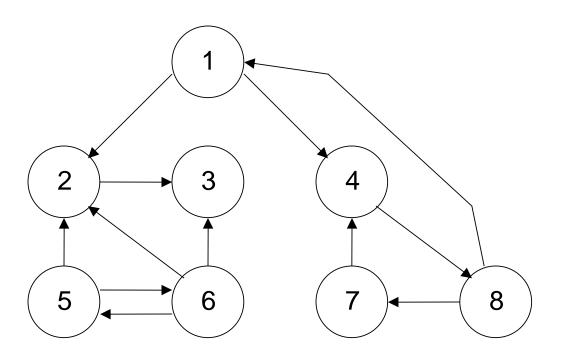
- The complete algorithm for finding the articulation points of a graph G is:
 - 1. Conduct a depth first traversal of G, producing the spanning tree T
 - 2.Traverse T in preorder and record the preorder sequence number prenum[v]
 - 3.Traverse T in postorder. For each node v calculate highest[v] as the minimum of
 - 1.prenum[v]
 - 2.prenum[w] for each node w such that G contains {v, w} and T does not.
 - 3.highest[x] for every child x of v
- 4. Determine the articulation points of G as follows
 - 1. The root is an articulation point if it has more then one child
 - 2.Any other node v is an articulation point if it has at least one child x such that highest[x] > prenum[v]

- The identification of articulation points is important in determine the critical components of networks.
- A component which corresponds to an articulation point is critical.
- If such a component fails, the network is compromised.

Depth-first Traversal: Directed Graphs

- The same algorithm can be used as for a non-directed graph.
- We only need to change the definition of adjacency to take account of the direction of each edge.
- The result of using this algorithm on a directed graph is typically quite different from its use on a nondirected graph.
- The result of the algorithm is typically a forest of sub-trees which together span the graph

– Consider the graph:



Traversing starting at node 1

```
•dfv(1) initial call
```

dfv(2) recursive call

dfv(3) recursive call; progress blocked

dfv(4) an unvisited neighbour of node 1

dfv(8) recursive call

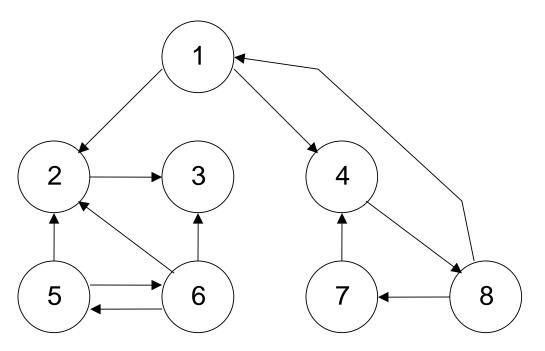
dfv(7) recursive call; progress blocked

dfv(5) new starting point

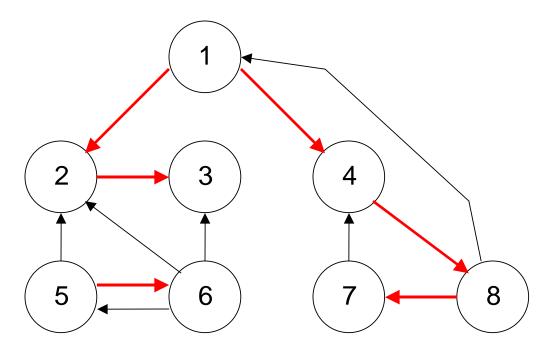
dfv(6) recursive call; progress blocked

There are no more nodes to visit

– Using the example graph:



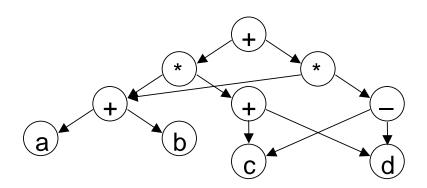
Using the example graph, the spanning forest F is:



- For a non-directed graph the set of edges of G not in T all connect nodes to their ancestors.
- —This is not the case for edges in the directed graph G which are not in F.
- –In this case, the edges fall into 3 groups:
 - Those like (3, 1) which connect a node to its ancestor.
 - Those like (1, 8) which connect a node to its descendant.
 - Those like (5, 2) which connect two unrelated nodes.

Acyclic graphs

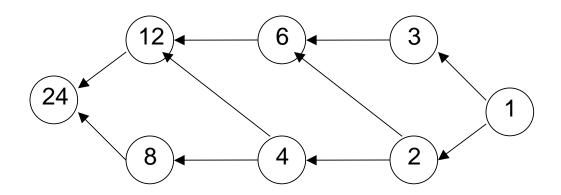
- Directed acyclic graphs can be used to represent several interesting data structures.
- For example representation of a arithmetic expression with a common sub expression.



$$(a + b)(c + d) + (a + b)(c - d)$$

Acyclic graphs

 Acyclic graphs offer a convenient way to represent partial orderings.

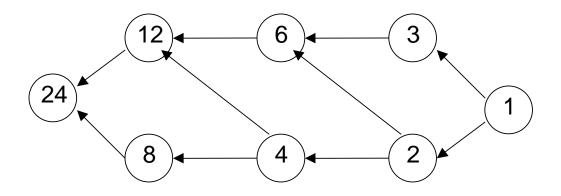


- ... is a factor of.

Acyclic graphs

- Another example of a partial ordering occurs in project management where the nodes represent milestones in the project and the edges represent activities.
- A topological ordering is one in which if (i, j) is an edge then i comes before j in the ordering of the nodes.
- Using the ... is a factor of graph as an example.

Acyclic graphsTopological sorting



- Valid topological sorts are:
 - •1, 2, 3, 4, 6, 8, 12, 24
 - •1, 3, 2, 4, 6, 8, 12, 24
 - •1, 2, 3, 6, 4, 8, 12, 24
 - •etc

Acyclic graphsTopological sorting

– The depth first visit algorithm can be easily modified to do a topological sort:

```
for each v ∈N do
    visited[v] = false
for each v ∈N do
    if not visited[v] then
        tops[v]

function tops[v]
  visited[v] = true
  for each node w adjacent to v do
    if not visited[w] then
        tops[w]
        write v
```

function topsort(G)